

**AMENDMENTS TO THE CLAIMS:**

This listing of claims will replace all prior versions, and listings, of claims in the application:

**Listing of Claims:**

**Claims 1-7 (Cancelled).**

**8. (Currently Amended)** An isolated variant of an *Erysipelothrix rhusiopathiae* surface protective antigen SpaA protein or of a shortened form thereof (known as ΔSpaA protein),

wherein the SpaA protein has comprises the amino acid sequence encoded by the nucleotide sequence of SEQ ID NO:7, and the ΔSpaA protein is a shortened form of the SpaA protein in which ~~207~~the C-terminal 206 amino acid residues ~~at the C-terminus~~ of the SpaA protein of the SE9 strain of *Erysipelothrix rhusiopathiae* are deleted,

wherein said variant is immunogenic, and expressed in *E. coli* as inclusion bodies, and is selected from the group consisting of:

(1) the SpaA protein ~~of the SE9 strain of~~ *Erysipelothrix rhusiopathiae* comprising the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution ~~at~~of arginine to glycine at the residue corresponding to residue position 531 (~~arginine to glycine~~)of SEQ ID NO:2;

- (2) the SpaA protein of the SE9 strain of *Erysipelothrix rhusiopathiae* comprising the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitutions at of histidine to glutamine at the residue corresponding to residue position 214 of SEQ ID NO:2 (histidine to glutamine) and at the amino acid substitution of methionine to threonine at the residue corresponding to residue position 253 of SEQ ID NO:2 (methionine to threonine);
- (3) the ΔSpaA protein of the SE9 strain of *Erysipelothrix rhusiopathiae* comprising the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitutions at of histidine to glutamine at the residue corresponding to residue position 214 of SEQ ID NO:2 (histidine to glutamine) and at the amino acid substitution of methionine to threonine at the residue corresponding to residue position 253 of SEQ ID NO:2 (methionine to threonine);
- (4) the ΔSpaA protein of the SE9 strain of *Erysipelothrix rhusiopathiae* comprising the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitutions at of glutamic acid to glycine at the residue corresponding to residue position 69 of SEQ ID NO:2 (glutamic acid to glycine), at the amino acid substitution of glutamic acid to glycine at the residue corresponding to residue position 154 of SEQ ID NO:2 (glutamic acid to glycine), and at

the amino acid substitution of isoleucine to threonine at the residue corresponding to residue position 203 of SEQ ID NO:2 (isoleucine to threonine); and

(5) the ΔSpaA protein of the SE9 strain of *Erysipelothrix rhusiopathiae* comprising the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution at of aspartic acid to glycine at the residue corresponding to residue position 278 of SEQ ID NO:2 (aspartic acid to glycine).

**Claims 9-16 (Cancelled).**

**17. (Currently Amended)** A composition comprising as an active ingredient an isolated variant of an *Erysipelothrix rhusiopathiae* surface protective antigen SpaA protein or of a shortened form thereof (known as ΔSpaA protein),

wherein the SpaA protein has comprises the amino acid sequence encoded by the nucleotide sequence of SEQ ID NO:7, and the ΔSpaA protein is a shortened form of the SpaA protein, in which ~~207~~the C-terminal 206 amino acid residues ~~at the C-terminus~~ of the SpaA protein of the SE9 strain of Erysipelothrix rhusiopathiae are deleted,

wherein said variant is immunogenic, and expressed in *E. coli* as inclusion bodies, and is selected from the group consisting of:

(1) the SpaA protein of the SE9 strain of *Erysipelothrix rhusiopathiae* comprising the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution at of arginine to glycine at the residue corresponding to residue position 531 (arginine to glycine) of SEQ ID NO:2;

(2) the SpaA protein of the SE9 strain of *Erysipelothrix rhusiopathiae* comprising the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitutions at of histidine to glutamine at the residue corresponding to residue position 214 of SEQ ID NO:2 (histidine to glutamine) and at the amino acid substitution of methionine to threonine at the residue corresponding to residue position 253 of SEQ ID NO:2 (methionine to threonine);

(3) the ΔSpaA protein of the SE9 strain of *Erysipelothrix rhusiopathiae* comprising the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitutions at of histidine to glutamine at the residue corresponding to residue position 214 of SEQ ID NO:2 (histidine to glutamine) and at the amino acid substitution of methionine to threonine at the residue corresponding to residue position 253 of SEQ ID NO:2 (methionine to threonine);

(4) the ΔSpaA protein of the SE9 strain of *Erysipelothrix rhusiopathiae* comprising the amino acid

sequence encoded by SEQ ID NO:7 with the amino acid substitutions at of glutamic acid to glycine at the residue corresponding to residue position 69 of SEQ ID NO:2 (glutamic acid to glycine), at the amino acid substitution of glutamic acid to glycine at the residue corresponding to residue position 154 of SEQ ID NO:2 (glutamic acid to glycine), and at the amino acid substitution of isoleucine to threonine at the residue corresponding to residue position 203 of SEQ ID NO:2 (isoleucine to threonine); and

(5) the ΔSpaA protein of the SEQ strain of *Erysipelothrix rhusiopathiae* comprising the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution at of aspartic acid to glycine at the residue corresponding to residue position 278 of SEQ ID NO:2 (aspartic acid to glycine).

**Claims 18-25 (Cancelled).**

**26. (Withdrawn)** A nucleic acid encoding the variant of claim 8.

**Claims 27-36 (Cancelled).**

**37. (Withdrawn)** A method for immunizing against infection with *Erysipelothrix rhusiopathiae*, comprising administering the variant of claim 8 to an animal in need of immunization.

**Claims 38-45 (Cancelled).**

**46. (New)** The isolated variant of claim 8, wherein the SpaA protein comprises the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution of arginine to glycine at the residue corresponding to residue position 531 of SEQ ID NO:2.

**47. (New)** The isolated variant of claim 8, wherein the SpaA protein comprises the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution of histidine to glutamine at the residue corresponding to residue position 214 of SEQ ID NO:2 and the amino acid substitution of methionine to threonine at the residue corresponding to residue position 253 of SEQ ID NO:2.

**48. (New)** The isolated variant of claim 8, wherein the ΔSpaA protein comprises the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution of histidine to glutamine at the residue corresponding to residue position 214 of SEQ ID NO:2 and the amino acid substitution of methionine to threonine at the residue corresponding to residue position 253 of SEQ ID NO:2.

**49. (New)** The isolated variant of claim 8, wherein the ΔSpaA protein comprises the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution of glutamic acid

to glycine at the residue corresponding to residue position 69 of SEQ ID NO:2, the amino acid substitution of glutamic acid to glycine at the residue corresponding to residue position 154 of SEQ ID NO:2, and the amino acid substitution of isoleucine to threonine at the residue corresponding to residue position 203 of SEQ ID NO:2.

**50. (New)** The isolated variant of claim 8, wherein the ΔSpaA protein comprising the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution of aspartic acid to glycine at the residue corresponding to residue position 278 of SEQ ID NO:2.

**51. (New)** The composition of claim 17, wherein the SpaA protein comprises the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution of arginine to glycine at the residue corresponding to residue position 531 of SEQ ID NO:2.

**52. (New)** The composition of claim 17, wherein the SpaA protein comprises the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution of histidine to glutamine at the residue corresponding to residue position 214 of SEQ ID NO:2 and the amino acid substitution of methionine to threonine at the residue corresponding to residue position 253 of SEQ ID NO:2.

**53. (New)** The composition of claim 17, wherein the ΔSpaA protein comprises the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution of histidine to glutamine at the residue corresponding to residue position 214 of SEQ ID NO:2 and the amino acid substitution of methionine to threonine at the residue corresponding to residue position 253 of SEQ ID NO:2.

**54. (New)** The composition of claim 17, wherein the ΔSpaA protein comprises the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution of glutamic acid to glycine at the residue corresponding to residue position 69 of SEQ ID NO:2, the amino acid substitution of glutamic acid to glycine at the residue corresponding to residue position 154 of SEQ ID NO:2, and the amino acid substitution of isoleucine to threonine at the residue corresponding to residue position 203 of SEQ ID NO:2.

**55. (New)** The composition of claim 17, wherein the ΔSpaA protein comprises the amino acid sequence encoded by SEQ ID NO:7 with the amino acid substitution of aspartic acid to glycine at the residue corresponding to residue position 278 of SEQ ID NO:2.